

Serial Number: 08/876/132

CRF Processing Date: 9/9/98
 Edited by: A
 Verified by: (STIC staff)

RECEIVED

APR 28 2000

ENTERED
 TECH CENTER 1600/2900

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included:
- ☐ Deleted extra, invalid, headings used by an applicant, specifically:
- ☒ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as
- ☐ Inserted mandatory headings, specifically:
- ☐ Corrected an obvious error in the response, specifically:
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically:
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected:
- ☐ Other:

*Examin r: Th above corrections must b communicated to th applicant in th first Office Action. DO NOT send a copy of this form.

INPUT SET: S28494.raw

SEQUENCE LISTING

(1) General Information

- (i) APPLICANT: Fowler, Timothy
Stuart, Causey
- (ii) TITLE OF THE INVENTION: ENTEROBACTERIACEAE FERMENTATION
STAINS
- (iii) NUMBER OF SEQUENCES: 3
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Genencor International, Inc.
 - (B) STREET: 925 Page Mill Road
 - (C) CITY: Palo Alto
 - (D) STATE: CA
 - (E) COUNTRY: US
 - (F) ZIP: 94304-1013
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Diskette
 - (B) COMPUTER: IBM Compatible
 - (C) OPERATING SYSTEM: DOS
 - (D) SOFTWARE: FastSEQ for Windows Version 2.0
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/876,132
 - (B) FILING DATE: 23-JUN-1997
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE:
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Glaister, Debra J.
 - (B) REGISTRATION NUMBER: 33,888
 - (C) REFERENCE/DOCKET NUMBER: GC372
- (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 650-846-7620
 - (B) TELEFAX: 650-845-6504

(2) INFORMATION FOR SEQ ID NO:1:

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/876,132DATE: 09/16/98
TIME: 10:17:15

INPUT SET: S28494.raw

47 (i) SEQUENCE CHARACTERISTICS:
48 (A) LENGTH: 1660 base pairs
49 (B) TYPE: nucleic acid
50 (C) STRANDEDNESS: single
51 (D) TOPOLOGY: linear
52
53 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
54
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56 TAAAAACAAAT GAGTGAAGAA GAAAAGAAAA AAATAAATAC ATATTTTGAG TTAGTAAAAAG 120
57 AGAAAAGAAAA AATAAAAAGAA GACCTCGGCT TAACAGTCGA AAAACCAGAA ATAATAAAAA 180
58 GAAAGAGACT GTGATTTTTA ATGGAATCG TGAGGAAAAG AAAATTTTAA TTTTCATTTT 240
59 CGAGGGATTA ATTTGTTGTA AGTTGATGAA AAATCTAGAT AAAAAATGCA GATCAAAAAAT 300
60 GTGTTGAATT TGACATTATT GAAATACGTA GTATATCAAT AATGGGGTT TGTCTATTTT 360
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69 CAAGAAGTGC CTTTCGTTTT TGTATTTGTG AGAGAAATCA GCAACTGAGA AAAGAAGCTG 900
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84 (2) INFORMATION FOR SEQ ID NO:2:
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86 (i) SEQUENCE CHARACTERISTICS:
87 (A) LENGTH: 1847 base pairs
88 (B) TYPE: nucleic acid
89 (C) STRANDEDNESS: single
90 (D) TOPOLOGY: linear
91
92
93 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
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97 AGAGGGTTTG ACGAGGTTTT TTCGAGCTAT CCAGGGTTTT TGGGTTTTTG GGGTTGGATC 180
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PAGE: 3

RAW SEQUENCE LISTING PATENT APPLICATION US/08/876,132

DATE: 09/16/98
TIME: 10:17:17

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111 AAAAAATATT TGGAATTTTT TAAAGGCGAT ACTTGCTACC GCACCTTTGC CATATTTAAA 1020
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113 GAGTAAAAAC AATAACTTAT ATATTTAATT CTGAATTATA TTTGACAGTG ATTATTTAAT 1140
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119 TCAGCTAAAA CTGAAAATGA TTTTAAAAAA TACTCGTTCT TTATTCATTC GATCAATTGG 1500
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124 CGTATGAAAA TCCTATTGAT GATCCAGATG ATGGCATAAG AAAATCACAA TTCGCACGAA 1800
125 GAAATGCCTA TGCTTTCCGC ATTAAAAAAA CAAGCAAAAA GAGATCT 1847

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(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 371 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

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138 1 5 10 15
139
140 Trp Lys Lys Tyr Tyr Lys Lys Arg Arg Ser Asp Met Asp Phe Lys
141 20 25 30
142
143 Ser Arg Lys Leu Thr Leu Asn Glu Lys Lys Asp Leu Glu Lys Ile Tyr
144 35 40 45
145
146 Ala Glu Ser Glu Leu Lys Ala Lys Lys Leu Gly Thr Gln Pro Gly Val
147 50 55 60
148
149 Val Leu Glu Met Thr Met Lys Glu Met Met Lys Asn Ile Asn Leu Asp
150 65 70 75 80
151
152 Val Asn Glu Glu Thr Ala Gly Gln Tyr Arg Lys Leu Phe Lys Asn Lys

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RAW SEQUENCE LISTING PATENT APPLICATION US/08/876,132

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TIME: 10:17:19

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159	115	120	125
160			
161	Ile Cys Glu Arg Ile Gln Gln Leu Arg Lys Glu Ala Asp Asn Ala Arg		
162	130	135	140
163			
164	Arg Val Lys Asp Phe Asp Thr Met Lys Ala Lys Thr Lys Glu Ala Phe		
165	145	150	155
166			
167	Glu Leu Ser Phe Val Phe Asp Lys Asp Phe Leu Ser Glu Asn Arg Ile		
168	165	170	175
169			
170	Gln Trp Asn Asp Ile Ser His Asn Lys Lys Asp Ser Ala Ser Lys Arg		
171	180	185	190
172			
173	Lys Thr Met Lys Glu Ala Asp Thr Met Asp Asp Ile Phe Lys Arg Leu		
174	195	200	205
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176	Lys Asn Asn Lys Ser Thr Tyr Asp Arg Tyr Ala Gly Phe Leu Ser Ile		
177	210	215	220
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180	225	230	235
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184	Ala Lys Val Gly Asn Asp Arg Gly Gln Ser Glu Arg Thr Leu His Phe		
185	260	265	270
186			
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188	275	280	285
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191	290	295	300
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194	305	310	315
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196	Tyr Thr Leu Arg His Arg Val Ala Ser Asp Leu Lys Ala Ser Gly Ala		
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199	Asp Asp Phe Thr Ile Ala Ala Xaa Leu Gly His Arg Val Thr Gln Ser		
200	340	345	350
201			
202	Gln Glu Leu Leu Arg Leu Cys Ser Phe Val Xaa Arg Trp Tyr Arg Cys		
203	355	360	365
204			
205	Asn Trp Cys		

PAGE: 5

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/876,132

DATE: 09/16/98
TIME: 10:17:20

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206	370
207	
208	
209	

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/876,132

DATE: 09/16/98
TIME: 10:17:20

INPUT SET: S28494.raw

Line

Error

Original Text

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/876,132DATE: 09/09/98
TIME: 15:03:05

INPUT SET: S28494.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

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46

SEQUENCE LISTING

(1) General Information

(i) APPLICANT: Fowler, Timothy
Stuart, Causey

(ii) TITLE OF THE INVENTION: ENTEROBACTERIACEAE FERMENTATION
STAINS

(iii) NUMBER OF SEQUENCES: 3

(iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: Genencor International, Inc.
(B) STREET: 925 Page Mill Road
(C) CITY: Palo Alto
(D) STATE: CA
(E) COUNTRY: US
(F) ZIP: 94304-1013

(v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Diskette
(B) COMPUTER: IBM Compatible
(C) OPERATING SYSTEM: DOS
(D) SOFTWARE: FastSEQ for Windows Version 2.0

(vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER: 08/876,132
(B) FILING DATE: 23-JUN-1997

(vii) PRIOR APPLICATION DATA:
(A) APPLICATION NUMBER:
(B) FILING DATE:

(viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: Glaister, Debra J.
(B) REGISTRATION NUMBER: 33,888
(C) REFERENCE/DOCKET NUMBER: GC372

(ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: 650-846-7620
(B) TELEFAX: 650-845-6504

(2) INFORMATION FOR SEQ ID NO:1:

Does Not Comply
Corrected Diskette Needed

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/876,132DATE: 09/09/98
TIME: 15:03:05

INPUT SET: S28494.raw

47 (i) SEQUENCE CHARACTERISTICS:
48 (A) LENGTH: 1660 base pairs
49 (B) TYPE: nucleic acid
50 (C) STRANDEDNESS: single
51 (D) TOPOLOGY: linear
52
53 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
54
55 AGATCTACAC AAGGCAAAATT GAAAAAATAG ATAAAAATTTT CGCAGGTATT AAAGCCGACT 60
56 TAAAAACAAAT GAGTGAAGAA GAAAAGAAAA AAATAAATAC ATATTTTGTAG TTAGTAAAAAG 120
57 AGAAAAGAAAA AATAAAAGAA GACCTCGGCT TAACAGTCGA AAAACCAGAA ATAATAAAAA 180
58 GAAAGAGACT GTGATTTTAA ATGGAATCG TGAGGAAAAAG AAAATTTTAA TTTTCATTTT 240
59 CGAGGGATTA ATTTGTTGTA AGTTGATGAA AAATCTAGAT AAAAAATGCA GATCAAAAAAT 300
60 GTGTTGAAAT TGACATTAT GAAATACGTA GTATATCAAT AATGGGGGTT TGTCATTTTT 360
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68 AATCAGATGA TCTAGTAACG GGACTATTAG AGTGTGGAAC TCGAAATAGT TTTGATAAAA 840
69 CAAGAAGTGC CTTTCGTTTT TGATTTTGTG AGAGAATTCA GCAACTGAGA AAAGAAGCTG 900
70 ATAATGCAAG AAGAGTAAAA GATTTTCGATA CAATGAAAGC AAAAACTAAA GAGGCTTTTTG 960
71 AATTGAGTTT TGTTTTTGAT AAGGATTTTT TGAGTGAAAA TAGAATTCAA TGGAATGATA 1020
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74 TCCTTCTAT TTGTTCGATT ACAGGTGCA GACCAGCAGA AGTTTTAAAG GGTATAGAGA 1200
75 TAGTAAGAAA CAGATATGAG GATGGTATAT CTTTTAAAAAT ACTTGGTGCA AAGGTTGGAA 1260
76 ATGACAGAGG GCAAAGCGAA AGAACATTAC ATTTTGATTT ATCAAAATAT CATGATAATG 1320
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80 TAGCGGCTNT TTTGGGTCAC AGAGTGACTC AAAGCCAGGA GTTACTACGG CTATGCTCGT 1560
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82 AAGAGTCAGT TNGCTGTATC AAGGACTCCG AGCCAGATCT 1660
83
84 (2) INFORMATION FOR SEQ ID NO:2:
85
86 (i) SEQUENCE CHARACTERISTICS:
87 (A) LENGTH: 1847 base pairs
88 (B) TYPE: nucleic acid
89 (C) STRANDEDNESS: single
90 (D) TOPOLOGY: linear
91
92
93 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
94
95 AGATCTCAAC CAGTTTAAAA TCGCACTTCA AGAAGTAAAA ATAGGGGCGG GCACCGGCTC 60
96 TTTTTTGGT GTTTTGTAG TTAGTGGATA TATCTGTAG CTACAGAGAA AAGCGATTTT 120
97 AGAGGGTTTG ACGAGGTTTT TTGAGCTAT CCAGGGTTTT TGGGTTTTTG GGGTTGGATC 180
98 AGAAAAGTCG TTCAAGATTA TTGACATAAA GACAGGAAGG TTTATAACAA GTACCAGATA 240
99 CGACAAAACC AGCTTTGCAG GCTGGCTTTG AAGGACTAAA AGAAGTGGGG ACTTCTTTGA 300

RAW SEQUENCE LISTING
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TIME: 15:03:06

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(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 371 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

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139																
140	Trp	Lys	Lys	Tyr	Tyr	Lys	Lys	Arg	Arg	Ser	Asp	Met	Asp	Phe	Lys	
141				20				25					30			
142																
143	Ser	Arg	Lys	Leu	Thr	Leu	Asn	Glu	Lys	Lys	Asp	Leu	Glu	Lys	Ile	Tyr
144				35				40					45			
145																
146	Ala	Glu	Ser	Glu	Leu	Lys	Ala	Lys	Lys	Leu	Gly	Thr	Gln	Pro	Gly	Val
147				50				55					60			
148																
149	Val	Leu	Glu	Met	Thr	Met	Lys	Glu	Met	Met	Lys	Asn	Ile	Asn	Leu	Asp
150				65				70				75			80	
151																
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RAW SEQUENCE LISTING PATENT APPLICATION US/08/876,132

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TIME: 15:03:09

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157			
158	Gly Thr Arg Asn Ser Phe Asp Lys Thr Arg Ser Ala Phe Arg Phe Cys		
159	115	120	125
160			
161	Ile Cys Glu Arg Ile Gln Gln Leu Arg Lys Glu Ala Asp Asn Ala Arg		
162	130	135	140
163			
164	Arg Val Lys Asp Phe Asp Thr Met Lys Ala Lys Thr Lys Glu Ala Phe		
165	145	150	155
166			
167	Glu Leu Ser Phe Val Phe Asp Lys Asp Phe Leu Ser Glu Asn Arg Ile		
168	165	170	175
169			
170	Gln Trp Asn Asp Ile Ser His Asn Lys Lys Asp Ser Ala Ser Lys Arg		
171	180	185	190
172			
173	Lys Thr Met Lys Glu Ala Asp Thr Met Asp Asp Ile Phe Lys Arg Leu		
174	195	200	205
175			
176	Lys Asn Asn Lys Ser Thr Tyr Asp Arg Tyr Ala Gly Phe Leu Ser Ile		
177	210	215	220
178			
179	Cys Ser Ile Thr Gly Cys Arg Pro Ala Glu Val Leu Lys Gly Ile Glu		
180	225	230	235
181			
182	Ile Val Arg Asn Arg Tyr Glu Asp Gly Ile Ser Phe Lys Ile Leu Gly		
183	245	250	255
184	Ala Lys Val Gly Asn Asp Arg Gly Gln Ser Glu Arg Thr Leu His Phe		
185	260	265	270
186			
187	Asp Leu Ser Lys Tyr His Asp Asn Glu Gln Met Asn Tyr Ile Leu Ser		
188	275	280	285
189			
190	Gln Leu Lys Asp Asn Lys Phe Phe Tyr Lys Pro Asp Gly Lys Leu Tyr		
191	290	295	300
192			
193	Asn Ser Leu Arg Gln Tyr Leu Tyr Ile Gln His Arg Thr Phe Ser Leu		
194	305	310	315
195			
196	Tyr Thr Leu Arg His Arg Val Ala Ser Asp Leu Lys Ala Ser Gly Ala		
197	325	330	335
198			
199	Asp Asp Phe Thr Ile Ala Ala Xaa Leu Gly His Arg Val Thr Gln Ser		
200	340	345	350
201			
202	Gln Glu Leu Leu Arg Leu Cys Ser Phe Val Xaa Arg Trp Tyr Arg Cys		
203	355	360	365
204			
205	Asn Trp Cys		

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RAW SEQUENCE LISTING
PATENT APPLICATION , US/08/876,132

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TIME: 15:03:10

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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION . US/08/876,132

DATE: 09/09/98
TIME: 15:03:10

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